STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (httm, EFS Submission User Manual - ePAVE)

2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450

3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



TWP

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RAW SEQUENCE LISTING DATE: 04/17/2006
PATENT APPLICATION: US/10/574,469 TIME: 12:29:11
```

Input Set: A:\SQ-AIA-113.ST25.txt
Output Set: N:\CRF4\04172006\J574469.raw

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3 <110> APPLICANT: Aoki, Hirofumi
             Kodama, Tatsuhiko
             Hasegawa, Kiyotaka
     5
             Kajiya, Kentaro
      6
             Ishimatsu, Yumiko
             Ogou, Masashi
             Yoshida, Seiichi
             Kishimoto, Jiro
     10
     11
             Moro, Osamu
    13. <130> TITLE OF INVENTION: Methods for Predicting Skin Stain Formation and for a
Screening
             Agent Capable of Inhibiting Skin Stain Formation by Means of
    14
             Promoted Genes in Stained Regions as an Indicator
     15
     17 <130> FILE REFERENCE: AIA-113-PCT
C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/574,469
C--> 19 <141> CURRENT FILING DATE: 2006-03-31
     19 <150> PRIOR APPLICATION NUMBER: JP 2003-343549
                                                          Dres Not Comply
     20 <151> PRIOR FILING DATE: 2003-10-01
                                                           Corrected Diskette Needed
     22 <150> PRIOR APPLICATION NUMBER: JP 2003-344786
    23 <151> PRIOR FILING DATE: 2003-10-02
     25 <160> NUMBER OF SEQ ID NOS: 7
     27 <170> SOFTWARE: PatentIn version 3.3
    29 <210> SEQ ID NO: 1
                                        See erron explanation on page 5,
    30 <211> LENGTH: 758
    31 <212> TYPE: DNA
    32 <213> ORGANISM: Mouse
    35 <220> FEATURE:
     38 <400>_SEQUENCE: 1
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    43 cagaagggag aagacateet gecaagteet tgaaaeteag gegetgetgt caeetatete
                                                                             180
                                                                             240
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    47 acaagctacc agtcaagtca tgggtggtgc ctggggctct cccacagata tagggcctcc
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     49 tecgeccaga tgaagegttg atgeccagat gtggagacae cagaageata cacactatgt
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                                                                             420
    51 tgccttgccc cttgccaatg agctgtgaca ctggaatgct tcacttcaga catcagggcg
     53 gatggattgc agaattccaa gtcctcattc caaaggtgtc accaaccttc agagtcacta
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                                                                             600
     57 cctgtgggag ctacagatcc agagactttc aagctgacta gagtgcagag aagcaagacc
                                                                             660
     59 tcagtgtgat cagccgagac tacagcatct tgggaaccct cagtcagccc caaaccccta
     61 acacttaacc actggtctcc aaaccaacac ctgtaacttc ctaatgaaat catcaggagg
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     66 <210> SEQ ID NO: 2
     67 <211> LENGTH: 2063
     68 <212> TYPE: DNA
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/574,469 TIME: 12:29:11

Output Set: N:\CRF4\04172006\J574469.raw

Input Set : A:\SQ-AIA-113.ST25.txt

DATE: 04/17/2006

69 <213> ORGANISM: Homo sapiens 71 <400> SEQUENCE: 2 72 ctctgcccat tgggaaacac ctctctatga ctctataaat gtccaaggtg gccccaaggg 60 74 aggacttctg cagcacagct cccttcccag gacgtgaaaa tctgccttct caccatgagg 120 76 cttctagtcc tttccagcct gctctgtatc ctgcttctct gcttctccat cttctccaca 180. 78 gaagggaaga ggcgtcctgc caaggcctgg tcaggcagga gaaccaggct ctgctgccac 240 300 80 cgagtcccta gccccaactc aacaaacctg aaaggacatc atgtgaggct ctgtaaacca 82 tgcaagcttg agccagagcc ccgcctttgg gtggtgcctg gggcactccc acaggtgtag 360 84 cacteceaaa geaagaetee agacagegga gaaceteatg cetggeacet gaggtaceea 420 86 gcagcctcct gtctcccctt tcagccttca cagcagtgag ctgcaatgtt ggagggcttc 480 88 atctcgggct gcaaggaccc tgggaaagtt ccagaactcc acgtccttgt ctcaattgtg 540 90 ccatcaactt tcagagctat catgagccaa cctcagcctt ccgagtagct gggattacag 600 92 gtgtgcgcca ccacacccgg ctaatttttg ctttttttt ttttgagaca gaatctcgct 660 720 94 ctgttgctca ggctggagtg gagtgcagtg gcacgatctc agctcactgc agcctctgcc 780 96 tecegggtte aggagattet cetgaeteaa ceteetgagt agetgggatt acaggtgeet 98 gccactatgc ctagctaatt tttgtatttt tagtggagac ggggttttgc cgtgttggcc 840 900 100 ggggtggtct cgaactcctg acctcaagtg atccacccgc atcggccccc caaagtgctg 960 102 ggattatagg cgtgaaccac cgcgcctgtc ccattgttgt gtaattttaa taattagttt - 1020 · 104 tttaagtach tgattttatg ggcanatttt tgtgggatga ttggagttaa tcaaataaag 106 ettgleatyt gegtagtieg gtaagataae ttetttaaar teatgittie tetgeetiga 1080 108 ggtagtgagg gaaagatett aateagtatt ttggtaatta aetgattgaa tteaageaaa 1140 1200 110 tgagacatca tgaacttcag tggttattga tatttcaggg tatatacctg aaatgcctag 112 aggatacaga tttctcattt cattctttgg tctttcattt ctctatatac agaaatgaaa .1260 1320 114 tgacacttct gggaggcagt agaagcagga agtcaatgaa ttgagtagag ggtcccattc 116 cctcaggctg tcattgatca gtgacaattt ataaaaacaa actgcaaagt ctgtggcaag 1380 1440 118 tggctgcctg cttcctagaa ggagcccatg aaggttaaac tctgtggtcg gtatttgcaa 1500 120 gcgccgggcg tggtggctca cgcctgtaat cctagcacct tgggaggcca aggcaggcgg 122 atcacccgag gtcaggagtt tgaggatttg caagcaaaag gtcctctcct gagtctttcc 1560 124 cagataccca gcagtgcaga ggctagctgt ggaaggttgc agtgggacag gaatgtattg 1620 1680 126 tatgccttgc cttacttgtc accattgaga tttccagaga aatgggcata acgtctctta 128 acaacaacag cagaaagcaa aatacattaa cttaaggttg acaacaaaag attatcaagt 1740 130 accatgtttt ccaaccaacc agttattcgt ggtaataata aaataaaggt gggaaaatgt -1800 1860 132 tataattttt aaggaaactg tgtactttaa aaatcttctt tatgaatatc caatgttact 1920 134 gtaatcctgc tecattaaat geageattgt tgteaggtge tgeetettge ttgggaacag 136 cattgggctt ttaaatgtct gcagaatctc tgcgttcgaa gggaattgag aatgaacttc 1980. 138 ctggtactgt aatgaaaata aggtctgctc aacacagtaa acgtttcctc tcttctttaa 2040 2063 140 aaaaaaaaaa aaaaaaaaa aaa 143 <210> SEO ID NO: 3 144 <211> LENGTH: 742 145 <212> TYPE: DNA 146 <213> ORGANISM: Rat 148 <400> SEQUENCE: 3 149 ttttttgtg ccactgette etetetagee gtgagaetee agetaettea etatgegaet 60 120 151 teteaceete teeggtttgt tetteatget gtteetetgt etetgegtte teteeteaga 153 agggagaaag cgtcctgcca agttcccgaa actcaggcct gctgtcatct atctcctaga 180 240 155 tocaaaccaa taactggaaa ggaaaccaca caagaccctg cagaccatgc agaaagctag 300 157 aatccaattc atggggtggt geetgggget eteccaeaga tatagggeet eeegaagetg 360 159 geetecaceg agatgaaaeg ttgatgteca gttatggaga caacettetg geecetacea 420 161 accttcatgg ccagaaagct gtgacaccag aatgtttcac ttcagacagc tgaaggatta

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/574,469

DATE: 04/17/2006 TIME: 12:29:11

Input Set : A:\SQ-AIA-113.ST25.txt
Output Set: N:\CRF4\04172006\J574469.raw

	163	cagaattcca agecetegtt ccaaaggtgc aaccaacett cagagtcaet atgatecagg	480
	165	gtcagcccac aagtcttcat ggctcctgca gagtaaaagt ccaagattcc atccctggga	540
	167	gctacagatt cagagacttc caagctgact ggcgaacaga gtagcaagac ttccttgtga	600
	169	tragatraga ttaragrate ttaggaacee traggacacee craaacecat agratttaat	660
	171	caacgggata tgaaccaact cctgtaactt cctaatgtaa tcaccaggag aacaccaaaa	720
	173	ataataaatc ataaatcaat gt	742
	176	<210> SEQ ID NO: 4	
	177	<211> LENGTH: 20	•
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	179	<213> ORGANISM: Artificial	
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	182	<223> OTHER INFORMATION: Sense Primer	
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		<213> ORGANISM: Artificial	
٠.		<220> FRATURE: " " " " " " " " " " " " " " " " " " "	
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		<400> SEQUENCE: 5	. 20
	197	gacaaggatg agaaaacacg	20
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RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 04/17/2006

PATENT APPLICATION: US/10/574,469

TIME: 12:29:12

Input Set : A:\SQ-AIA-113.ST25.txt Output Set: N:\CRF4\04172006\J574469.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 6 V

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:4,5,6,7

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/574,469

DATE: 04/17/2006 TIME: 12:29:12

Input Set : A:\SQ-AIA-113.ST25.txt

Output Set: N:\CRF4\04172006\J574469.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which

residue n or Xaa represents.

Seq#:1; N Pos. 6

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/574,469

DATE: 04/17/2006 TIME: 12:29:12

Input Set : A:\SQ-AIA-113.ST25.txt
Output Set: N:\CRF4\04172006\J574469.raw

L:19 M:270 C: Current Application Number differs, Replaced Current Application No

L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:39 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:1

L:39 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:1 L:39 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:1

L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0